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PRODUCT INSERT

LIFECODES® HLA-SSO TYPING KITS

For RESEARCH USE ONLY, NOT FOR USE IN DIAGNOSTIC PROCEDURES

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DEFINITION OF SYMBOLS (Product Labels and Supplemental Documents)

Batch Code



Catalog Number



Temperature limitations



Upper limit of temperature



Use By Date



Keep away from light



Sufficient for N Tests



Do Not Freeze



Caution – See instructions for Use



Consult instructions for use



Manufacturer



Danger




KITS/REAGENTS BY CATALOG NUMBER

LCT-DQAB LIFECODES HLA-DQA1/B1 SSO Typing Kit, Catalog # 628930R

LC-DQAB LIFECODES HLA-DQA1/B1 Typing Kit for use with Luminex


Product # 200210-50R

Reagent	Product Number	Fill volume	Storage	 Sufficient for 50 samples
MX-DQAB LIFECODES HLA DQA1/B1 Master Mix	200200	870 µL	2 to 8°C	
BM-DQAB LIFECODES HLA DQA1/B1 Probe Mix ‡	200201	810 µL	2 to 8°C Protect From Light	
DS Dilution Solution	628515	19.7 mL	18 to 30°C	
TAQ LIFECODES Taq Polymerase	628076	25 µL	-10°C to -30°C	General Purpose Reagent

LCT-DPAB LIFECODES HLA-DPA1/B1 SSO Typing Kit, Catalog # 628936R

LC-DPAB LIFECODES HLA-DPA1/B1 Typing Kit for use with Luminex


Product # 200110-50R

Reagent	Product Number	Fill volume	Storage	 Sufficient for 50 samples
MX-DPAB LIFECODES HLA DPA1/B1 Master Mix	200100	870 µL	2 to 8°C	
BM-DPAB LIFECODES HLA DPA1/B1 Probe Mix	200101	810 µL	2 to 8°C Protect From Light	
DS Dilution Solution	628515	19.7 mL	18 to 30°C	
TAQ LIFECODES Taq Polymerase	628076	25 µL	-10°C to -30°C	General Purpose Reagent

LCT-DQA LIFECODES HLA-DQA1 SSO Typing Kit, Catalog # 628933

LM-DQA LIFECODES HLA-DQA Typing Kit for use with Luminex


Product # 628061R

Reagent	Product Number	Fill volume	Storage	 Sufficient for 20 samples
MX-DQA LIFECODES HLA DQA Master Mix	628062	420 µL	2 to 8°C	
BM-DQA LIFECODES HLA DQA Probe Mix ‡	628063	360 µL	2 to 8°C Protect From Light	
DS Dilution Solution	628155	9.9 mL	18 to 30°C	
TAQ LIFECODES Taq Polymerase	628076	25 µL	-10°C to -30°C	General Purpose Reagent

LCT-DPB LIFECODES HLA-DPB1 SSO Typing Kit, Catalog # 628937

LM-DPB LIFECODES HLA-DPB Typing Kit for use with Luminex

Product # 628910 R

Reagent	Product Number	Fill volume	Storage	 Sufficient for 20 samples
MX-DPB LIFECODES HLA DPB Master Mix	628903	420 µL	2 to 8°C	
BM-DPB LIFECODES HLA DPB Probe Mix ‡	628902	360 µL	2 to 8°C Protect From Light	
DS Dilution Solution	628155	9.9 mL	18 to 30°C	
TAQ LIFECODES Taq Polymerase	628076	25 µL	-10°C to -30°C	General Purpose Reagent

‡Probe Mixes are light sensitive: keep exposure to light at a minimum.
CAUTION: Do not use components past their expiration dates.

SUMMARY AND EXPLANATION

DNA-based HLA typing using PCR amplified DNA is a common laboratory procedure. PCR amplification of DNA is used as the means to enrich for a selected DNA region. For HLA typing, a subsequent assay is utilized to determine the properties of the amplified DNA. Several types of assays, such as SSP (1), direct SSOP (2), RFLP (3) and reverse SSOP dot blot technologies (4), have been used in HLA typing. Like SSOP and reverse dot blot methods, LIFECODES HLA-SSO Typing kits utilize sequence-specific oligonucleotides (SSOs) to identify which HLA alleles are present in a PCR amplified sample. It is the set of SSOs employed, not the methodologies that determines the ability to distinguish among the various alleles present in the PCR amplification. Whereas reverse dot blot and SSOP methods employ enzyme labels and colorimetric substrates that require subsequent development, the LIFECODES assay is a homogenous multiplex system. That is, all SSOs are analyzed simultaneously and the entire assay is carried out in a single reaction vessel with the addition of a single reagent.

PRINCIPLES OF THE PROCEDURE

The LIFECODES HLA-SSO Typing procedure is based on the hybridization of labeled single stranded PCR product to SSO probes. Amplification of DNA using PCR typically employs equimolar amounts of both forward and reverse primer to generate a double-stranded DNA product. However, if the amount of one primer is in excess relative to the other, the reaction will generate some single-stranded DNA product in addition to double-stranded product. During the initial cycles of the LIFECODES amplification step, double-stranded DNA is generated. Once the limiting primer is exhausted, the remaining primer uses the double-stranded product as a template for generation of single-stranded DNA. This method generates both double stranded and single stranded products that upon denaturation, will both participate in the hybridization reaction.

Each of the different probes may be homologous to a sequence within the amplified DNA that is unique to an allele or group of alleles. In other words, these probes are designed so that each probe preferentially hybridizes to a complementary region that may or may not be present in the amplified DNA. In addition, the amplified DNA is also hybridized to one or more Consensus probes homologous to sequences present in all the alleles of a locus. SSO Typing can be affected by the type of biological material, method of purification, amount and integrity of genomic DNA. Therefore, the signal obtained for the Consensus probe(s) can serve as an indicator of the success of the amplification and hybridization procedures. Also, the signal obtained with the Consensus probe can be used to normalize the signal of the allele specific probes and correct for variations in the amount of amplified product in the hybridization reaction. The analysis of the results generated from the SSO typing can be used to determine the presence or absence of particular DNA sequences in amplified DNA and to identify the possible alleles in the sample.

For the LIFECODES HLA-SSO Typing procedure, probes are attached to Luminex Microspheres designed for use with the Luminex Instrument. Up to 100 different populations of Luminex Microspheres can be mixed together and analyzed by the Luminex Instrument because each population of microspheres can be distinguished by its unique fluorescence signature or color. A different SSO probe can be attached to each color microsphere. Therefore, a mixture of several probes can be distinguished from each other by virtue of their association with particular color microspheres. The Luminex Instrument is also able to quantify the relative amounts of labeled PCR product hybridizing to each Luminex Microsphere. Therefore, the relative signal obtained with the SSO probes in the LIFECODES assay, as with other SSOP methods, can be used to assign the probes as having positive or negative reactivity with the amplified DNA sample (see Results section). This in turn provides the information needed to determine the HLA phenotype of the sample.

REAGENTS

A. Identification

See tables in Reagents by Catalog Number section for complete listing of products and catalog numbers.

B. Warnings or Cautions

1. For Research Use Only. Not for use in diagnostic procedures.
2. Separate pipettes should be designated for Pre-PCR manipulations as well as for Post-PCR manipulations.
3. **Biohazard:** All biological and blood samples should be treated as potentially infectious. **Use Universal Precautions when handling.**
4. Dilution Solution, Probe mixes, Taq Polymerase and R-Phycoerythrin Conjugated Streptavidin contain hazardous compounds. Avoid contact with skin and eyes and dispose of all materials after use according to local regulations. See Material Safety Data Sheets for additional information.

C. Storage Instructions

1. Refer to kit component packaging label for proper storage temperatures.
2. Probe mixes and R-Phycoerythrin Conjugated Streptavidin are light sensitive, **KEEP FROM LIGHT; DO NOT FREEZE.**
3. Do not use components past their expiration date.

D. Purification or Treatment Required for Use

See "Specimen Collection and Preparation."

E. Instability Indications

1. If salts have precipitated out of solution during shipping or storage, resolubilize completely prior to use by vortexing at room temperature (18 to 30°C).
2. Do not use R-Phycoerythrin Conjugated Streptavidin that has been frozen during shipment or storage.

INSTRUMENT REQUIREMENTS

1. Luminex Instrument and XY Platform (Product Number 888300, 888310)
2. The following Thermal Cyclers have been validated: 96-Well GeneAmp® PCR System 9700 set to MAX mode (Base Cat # N8050200, Gold Block Cat #4314878), Veriti™ 96-Well Thermal Cycler set to 9700 MAX mode (Cat #4375786). Refer to Table 2 for maximum ramp speeds.

SPECIMEN COLLECTION AND PREPARATION

- a. Human DNA can be purified from Whole blood, Buffy coats and Buccal swabs using a validated method that meets the criteria below.
- b. DNA extracted from blood preserved in EDTA and ACD (Acid Citrate Dextrose) have been tested and shown to yield expected performance in this assay. DNA extracted from blood preserved in heparin cannot be used in this assay. Other preservatives have not been tested.
- c. The isolated DNA should be in 10 mM TRIS, pH 8.0-9.0, or in nuclease free water. If a chelating agent such as EDTA is present the final concentration of the chelating agent should not exceed 0.5 mM.
- d. The presence of alcohol, detergents or salts may adversely affect DNA amplification.
- e. Final DNA concentration should be 10 to 200 ng/μL.
- f. Absorbance measurements of the DNA sample at 260 and 280nm should give a ratio of 1.65 to 2.0.
- g. DNA can be used immediately after isolation or stored at -20°C for up to 1 year. Repeated freeze/thawing should be avoided since this can result in DNA degradation.

PROCEDURE

A. Materials Provided (See tables in Reagents by Catalog Number section for specific information)

- Appropriate Master Mix (MX)
- Appropriate Probe Mix (BM)
- Dilution Solution (DS)
- Threshold Table(s), Probe Hit Chart(s)
- LIFECODES Taq Polymerase* (LIFECODES Cat. No 628076)

B. Materials, Required, but Not Provided

The following are recommended for use with the kit:

- Luminex Sheath Fluid (1x LIFECODES Cat. No. 628005)
- Nuclease-free water (LIFECODES Cat. No. 757003; 20mL)
- PCR tubes and caps - Corning® Thermowell Tube Strips (Costar® Cat. No. 6542, LIFECODES Cat. No. 888640) or Corning® Thermowell PCR 96 well plates (Cat. No. CLS6551) or ThermoFisher AB Gene® Superplate 96-well PCR plate (Cat. No. AB-2100)
- Costar® plate (Costar® Cat. No. 6509, LIFECODES Cat. No. 888630)
- Thermowell Clear Polyethylene Tape (Costar® No. 6524 (LIFECODES Cat. No. 888635)
- R-Phycoerythrin Conjugated Streptavidin (SA-PE), 1mg/mL (LIFECODES Cat No. 628511)
- Luminex Calibration Kits (Luminex 100/200 Calibration Kit, Luminex 100/200 Performance Verification Kit, LIFECODES Cat. Nos. 628018 and 628019 respectively)

C. Additional Materials to be provided by the user

- Vortex Mixer
- Silicone compression Mat. Axygen Scientific , # CM-FLAT, or equivalent
- Bath Sonicator
- Microcentrifuge Barrier filter tips
- Pipettors, Multichannel pipettors and tips (1-20µL, 20-200µL, 1000µL)
- Spreadsheet analysis software
- Heat Block
- 70% Isopropanol or 20% Bleach
- Retainer tray – Applied Biosystems #403081 (for use with the 9700 thermal cycler only)

DIRECTIONS FOR USE

NOTES:

- Probe mixes and SA-PE are light sensitive: **keep away from light and do not freeze.**
- Warm beads at 55° to 60°C for at least 5 -10 minutes to thoroughly solubilize components in probe mixture.
- Sonicate briefly (~15 sec), then vortex probe mix for about 15 seconds to thoroughly suspend the beads.
- Take extreme caution in the aliquoting process, using calibrated pipettes. Failure to do so may result in reagent loss and sample failure.
- All temperatures must be precisely maintained. Fluctuations as little as +/- 0.5°C can affect results.
- At the hybridization stage, samples should not remain in the diluted state at 56°C for more than 5 minutes (see Results section).
- It is recommended to assay the amplified samples as soon as possible. If the samples cannot be run on the Luminex Instrument the same day, ***the amplified product can be stored up to 3 days at 2-8°C prior to use. For longer storage, store at -20°C up to one week until ready to assay. The amplified product can only be frozen and thawed once. Repeated freezing and thawing will result in degradation of amplified samples and will yield poor results if assayed.***

A. Purify genomic DNA, using method of choice; final concentration should be 10 to 200 ng/µL. Adjust, if necessary, with nuclease free water. Keep all samples at similar concentrations.

B. DNA amplification (PCR)

1. Allow the Master Mix to warm to room temperature (18 to 30°C).
2. Gently vortex for approximately 10 seconds. This will ensure the salts are in solution. Spin briefly (5 – 10 seconds) in microcentrifuge to bring contents to the bottom of the tube.
3. Using **Table 1** below, prepare the components for amplification for n+1 reactions using the indicated amount of each component per reaction (except for DNA). Bring to a final volume of 20µL per reaction with nuclease free water. Gently vortex.
4. Pipette the appropriate amount of Genomic DNA (**40 to 120ng**) into the PCR tubes.
5. Aliquot the amplification mix into the PCR tubes containing the genomic DNA. (The total volume of amplification mix and genomic DNA should equal 20µL for each sample reaction.)
6. Cap tubes tightly to prevent evaporation during PCR.
7. Place samples in the thermal cycler and run program, see **Table 2 and 3**.

Table 1. Reaction Components for Amplification

Component	Amount per PCR sample reaction
LIFECODES Master Mix	6µL
Genomic DNA 10-200ng/µL	Total of ~80ng
LIFECODES Taq Polymerase	0.2µL (1U)
Nuclease-free water	To 20µL final volume

Table 2. Thermal Cycler Conditions for Amplification

Thermal Cycler	Mode (Ramp speed)
GeneAmp® PCR System 9700	MAX mode (3.9°C/sec)
Veriti™ 96-Well Thermal Cycler	9700 MAX mode (3.9°C/sec)

Table 3. Thermal Cycler Conditions for Amplification

Step	Temperature and Incubation Time	# of Cycles
1	95° C for 3 min	1
2	95° C for 15 sec	12
	60° C for 30 sec	
	72° C for 30 sec	
3	95° C for 10 sec	28
	63° C for 30 sec	
	72° C for 30 sec	
4	72° C for 2 min	1
5	4° C forever	1

Note: To be sure of sample amplification, refer to Product Gel Electrophoresis (Appendix A).

C. Hybridization

- Be sure hybridization buffer components of the LIFECODES probe mix are solubilized and that the beads are thoroughly suspended.
- Turn on the Luminex Instrument and XY Platform to allow for 30 minute warm-up.
 - Warm probe mix in a 55° to 60°C heat block for at least 5 -10 minutes to thoroughly solubilize components in probe mixture.
 - Sonicate briefly (~15 sec), then vortex probe mix for about 15 seconds to thoroughly suspend the beads.
 - Combine 15 µL of the **appropriate** probe mix with 5 µL of locus specific PCR product into each well of a thermal cycler 96 well plate (Costar® No. 6509). When aliquoting probe mix to more than 10 wells, gently vortex probe mix after each set of ten. Seal plate with polyethylene tape (Costar® No. 6524).
 - Place silicone compression mat on top of plate prior to hybridization.
 - Hybridize samples under the following incubation conditions:

Table 4. Thermal Cycler Conditions For Hybridization

97°C for 2 minutes
47°C for 10 minutes
56°C for 8 minutes
56°C HOLD

- Ensure that the detection laser on the Luminex Instrument is turned on at least 30 minutes before the hybridization ends.
- While the samples are hybridizing, prepare a 200:1 Dilution Solution/SA-PE mixture. Combine 170µL Dilution Solution (DS) and 0.85µL 1mg/mL SA-PE per sample. *It is recommended to make enough Dilution Solution Mixture for n+1 samples to account for pipetting loss. (See Table 5).*
- Keep Dilution Solution/ SA-PE mixture in the dark, at room temperature; SA-PE is light sensitive! The Dilution Solution may be warmed at 45°C for 5 minutes and vortexed upon arrival to ensure all components are in solution. Dilution solution must be at room temperature (18 to 30°C) before making the mixture. **Prepare prior to use and discard any remaining portion.**

Table 5. Dilution Solution Preparation Volumes

# of Samples	Dilution Solution (DS)	SA-PE
1	170µL	0.85µL
5	850µL	4.25µL
10	1700µL	8.5µL
20	3400µL	17µL
50	8500µL	42.5µL

Note: DO NOT CANCEL HYBRIDIZATION PROGRAM BEFORE REMOVING THE TRAY FROM THE THERMAL CYCLER!

8. At the 56°C hold, while the tray is on the thermal cyclor, dilute each sample with 170µL of the prepared Dilution Solution/ SA-PE mixture. **It is critical to dilute all samples within 5 minutes (following the 8 minute 56°C HOLD step).**
9. Remove the sample tray from the thermal cyclor and place in the Luminex Instrument.

D. Analyze sample using the Luminex Instrument*

For best results, assay the samples immediately using the Luminex Instrument. Samples can be read up to 30 minutes after being diluted. If not read immediately, protect samples from light.

1. Turn on the Luminex Instrument between 30 minutes and 4 hours before assaying the samples.
2. Prior to analyzing the samples on the Luminex Instrument, set up a Batch Run by which the samples will be analyzed.
 - a) Select **Create a New Batch** from the File menu.
 - For example, if analyzing for HLA-DPB1, add Batch for **HLA-DPB1**.
 - The Batch Template is provided and is named, in this case, HLA-DPB1.
 - Please note that the template versions are lot number specific and correspond to the kit lot numbers.
 - Follow the stepwise instructions that appear on the screen for creating batches.
 - **When naming the batch, do not include commas in the name because information after a comma will be lost upon exportation of the data.**
 - For further instructions on creating batches and multibatches, refer to the Luminex User's Manual
 - b) Click the eject icon to eject the plate holder. Place the 96 well thermal cyclor plate containing the samples in the XYP heater block present on the plate holder.
 - c) Click the Retract icon. The samples are now ready to be analyzed. A prime step should be performed before starting the run.
 - d) After the samples have been run through the instrument, a sanitization step with 70% Isopropanol or 20% household bleach should be performed followed by two wash steps. The instrument can be turned off at this point if it is not going to be used for the remainder of the day.
3. After a batch is complete, the data is exported as a comma separated values (csv) file. These files are named 'OUTPUT.CSV' and saved in a folder with the Batch Name. This data is then available for making typing assignments as described below.

*Refer to Luminex User's Manual for instrument operation, including daily startup, calibration, maintenance, and shutdown procedures.

RESULTS

Sample typing can be done as follows:

The generated CSV files can be opened and the data processed with common spreadsheet programs such as Microsoft Excel, Lotus 123, Corel Quattro Pro, or similar software. Analysis is comprised of the following steps:

- 1) Verify that the Number of Events for each SSO in each sample is at least 60. This information is found in the **DataType: Count** section of the CSV file.
- 2) Determine that the values for the Consensus probes for each sample are above their minimum Median Fluorescent Intensity or MFI. The minimum thresholds are lot specific and can be found in the Threshold Table.

Caution:

- To obtain reliable results, there must be sufficient data gathered by the Luminex Instrument.
 - Collect at least 60 events for HLA-DQA1 (Part # 628933) and HLA-DPB1 (Part # 628937).
 - Collect at least 40 events for HLA-DQA1/B1 (Part # 628930R) and HLA-DPA1/B1 (Part # 628936R).
- 3) Subtract the Background Control value for each probe from the sample values producing the background corrected data set. Background Control values are found in the Threshold Table and are lot specific. Background values are average MFI values for each bead to compensate for background noise due to bead variation.
 - 4) For each sample, divide the background-corrected data for each probe by the background-corrected value for the corresponding consensus probe producing the normalized data set.

$$\frac{\text{MFI (Probe)} - \text{MFI (Control blank for probe)}}{\text{MFI (Consensus)} - \text{MFI (Control blank for consensus)}}$$

- 5) For each probe, record the normalized value on the Threshold Table Worksheet.
- 6) Once all values have been assigned, the probe hit pattern (i.e., the combination of all positive and negative assignments for a given sample) can be compared with the Probe Hit Table(s) provided.

Caution:

- There is a separate threshold table for each locus and each probe mix.
- **These threshold tables are Lot-specific; be certain that the Lot # on the threshold tables matches the Lot # in the typing kit.**

- If a normalized value for a particular probe falls above the maximum threshold for a negative assignment and below the minimum value for a positive assignment, the sample should be considered as indeterminate for this probe. The sample should be typed, first assuming the value to be negative and then again assuming the value to be positive.
- See **EXPECTED VALUES** section for further information on threshold values.

QUALITY CONTROL

It is recommended that one negative and positive control be run with each test, such as a water blank and a previously typed sample respectively. Consensus SSO probes, listed on the Threshold Table, hybridize to their respective locus specific alleles. Values obtained with the Consensus SSOs from positive controls should exceed the threshold value for the SSO as set forth in the Threshold Table Worksheet. Values obtained with the Consensus SSOs from water blanks should be below the threshold value for the SSO as set forth in the Threshold Table Worksheet.

The LIFECODES Probe Mix(es) contain one or more consensus SSO probes identified in the typing kit worksheets. These consensus probes hybridize to all alleles and act as internal controls to verify amplification and to confirm that hybridizations occurred. If the minimum value is not obtained for these SSOs, the sample may not produce the correct typing and the sample test should be repeated.

The assay should be run as recommended in the package insert as well as performed with any other quality control procedures that are in accordance with local, state, federal and/or accreditation agencies requirements.

LIMITATIONS OF THE PROCEDURE

The PCR conditions and assay conditions described require precisely controlled conditions. Deviations from these parameters may lead to product failure.

All instruments must be calibrated according to manufacturer's recommendations and operated within manufacture's prescribed parameters.

- 1) Beads must be pre-warmed and well suspended prior to use. This ensures that the hybridization buffer components are in solution.
- 2) 47°C and 56°C incubations require a high degree of accuracy (+/- 0.5°C). A thermal cycler should be employed. Temperature should be verified, within wells of the 96 well thermal cycler plate, using a thermocouple (e.g., Bio-Rad, Model VPT-0300 or equivalent). The temperature within wells and among wells should not vary more than +/- 0.5°C.
- 3) Time at 56°C is critical and should not exceed a total of 13 minutes. This includes the 8-minute incubation plus no more than 5 minutes to dilute all the samples with Dilution Solution/ SA-PE mixture.
- 4) Once diluted, the samples are stable at room temperature (18 to 30°C) for up to 2 hours (protect from light). Since a full 96-well plate can take up to 1.5 hrs. To run through the Luminex Instrument, the analysis should be started no more than 30 minutes after dilution to ensure that the last sample is analyzed within the 2 hr. limit.
- 5) Do not mix components from other kits and lots.

Due to the complex nature of HLA typing, qualified personnel should review data interpretation and typing assignments.

TROUBLESHOOTING

PROBLEM	POSSIBLE CAUSE		SOLUTION
Low Bead Count	Probe Mix not well suspended		Prewarm, sonicate and vortex probe mix and repeat assay.
	Instrument not functioning properly	Out of Calibration	Calibrate Instrument. (Refer to Luminex IS User's manual.)
		Sample flow path blocked	Remove and sonicate needle. Perform backflush. Call Immucor Transplant Diagnostics, Inc. if problem persists. (888) 329-0255
CON Threshold Failure	Sample failed to amplify or amplified poorly*	Low DNA	Check DNA concentration and purity.
		Salts in Master Mix are out of solution	Heat Master Mix at 37°C for 5 minutes, vortex gently and spin down briefly.
		Poor Taq Polymerase	Use LIFECODES Taq Polymerase Catalog # 628076.
	Amplification conditions not within specific parameters		Run Thermal profile on Thermal cycler to verify parameters are within specified parameters.
	Low Median Fluorescent Intensity Value (MFI)		Warm dilution solution at 45°C for 5 minutes before use and vortex. Store at room temperature. Replace R-Phycoerythrin Conjugated Streptavidin.
Multiple SSO failures or sample fails to yield a HLA typing result	Allele specific amplification	Amplification conditions not within specific parameters	Run Thermal profile on Thermal cycler to verify parameters are within specified parameters.
	DNA sample contaminated		
	DNA partially degraded	Re-isolate DNA from Blood sample.	
	Evaporation during hybridization step		If not using an entire plate, leave one row empty on each side of samples to be assayed to allow plate to be sealed tightly.

* PCR amplification can be verified by gel electrophoresis (See Appendix A).

EXPECTED VALUES

Values typically can be resolved as positive and negative. In some rare instances, the values may be indeterminate. An "indeterminate value" represents a range in which neither positive nor negative values have been observed. If a sample contains indeterminate values for a particular SSO probe, the sample should be typed with the probe as a negative and again with the probe as a positive. Three cases can arise:

1. One of the choices (i.e. Positive or Negative) provides a match.
2. Both choices provide matches.
 - Sample can be re-assayed, or the alleles obtained from both typings can be reported.
3. Neither choice produces a match.
 - In this instance, there are likely to be other probes that were incorrectly assigned. This sample needs to be re-assayed and possibly re-amplified.
 - If more than two probes are indeterminate, the sample should be re-assayed.
 - If a probe hit pattern does not produce an HLA type, the sample should be re-amplified and re-assayed. It may also be necessary to re-isolate DNA from the sample and reamplify and reassay.
 - As noted in the **Limitations of the Procedure** section, it is critical to precisely follow the protocol. Any deviations can lead to sample typing failure.

REFERENCES

1. Olerup, O., et al. (1992) Tissue Antigens 39:225
2. Saiki, RK., et al. (1986) Nature 324: 163
3. Maeda, M., et al. (1989) Tissue Antigens 34: 290
4. Bugawan, TL., et al. (1990) Immunogenetics 32: 231

LIMITED LICENSES

Taq polymerase is manufactured for Immucor Transplant Diagnostics by Promega Corp. It is licensed to Promega under U.S. Patent Nos. 5,338,671 and 5,587,287 and their corresponding foreign patents. The purchase of this product includes a limited, non-transferable license under U.S. patent 5,981,180 or its foreign counterparts, owned by Luminex Corporation,

MANUFACTURER INFORMATION:

Manufacturer: Immucor Transplant Diagnostics, Inc., 550 West Avenue, Stamford, CT 06902. Phone: 203-328-9500, 888-329-0255
Fax: 203-328-9599

European Technical Service: Phone: +32/3 385 4791

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TRADEMARKS USED

AB Gene®	AB Gene House	Luminex®	Luminex Corporation
Costar®	Corning Incorporated	Gene Amp®	Roche Molecular System
Microseal™	Bio-Rad Laboratories, Inc.	Veriti™	Applied Biosystem
IDNA™ Agarose	Lonza Group, Ltd.	LIFECODES®	Immucor Inc.
GelStar™	Lonza Group, Ltd.		

Gel Electrophoresis

The PCR reactions performed in the LIFECODES HLA-SSO Typing Kits are designed to produce both double and single stranded products, which are the predominant products that hybridize to the SSOs. For quality assurance or to trouble shoot an experiment it might be necessary to perform gel electrophoresis to examine the PCR reaction for the presence of amplified DNA.

Materials Required (as listed or equivalent)

- Electrophoresis Grade Agarose (Lonza Group, Ltd. IDNA® Agarose No. 50170)
 - Electrophoresis apparatus/power supply
 - 1X Gel Buffer (40xTAE, Promega No. V4281)
- GelStar® Nucleic Acid Gel Stain (Lonza Group, Ltd. No.50535)
 - UV Transilluminator (ChromatoVUE, UVP Inc. Model TM36)
 - Photographic imaging system

The relative migration of the single stranded product is dependent upon the gel concentration and buffer system employed. Approximate migrations for each amplification are listed below for samples run in a 2% Agarose gel in 1X TAE buffer.

Electrophoresis Conditions

Note: Not applicable for HLA-DQA1/B1 (Part # 628930R) and HLA-DPA1/B1 (Part # 628936R) as individual bands cannot be distinguished in combination multiplex.

1. Remove GelStar® Nucleic Acid Stain (Lonza Group, Ltd. No.50535) from freezer to thaw. Keep in dark.

2. The gel used for this procedure must be 2%, i.e. for a 200ml gel bed use 4 grams of agarose to 200mL 1X TAE (Dilute from 40X TAE). Add 10µL GelStar® Nucleic Acid Stain to the molten agarose. When pouring the gel be sure to leave ample room for DNA to run a significant distance (1 to 2 inches). **USE CAUTION: GelStar® is a potential Carcinogen.**

NOTE: It is possible to run gels with 20µL of 10mg/mL Ethidium Bromide in place of GelStar® Nucleic Acid Stain. Product band intensity will be less in gels containing Ethidium Bromide than in gels containing GelStar®. **USE CAUTION: Ethidium Bromide is a known Carcinogen.**

3. Keep gel in dark and allow to solidify.

4. Load a mixture of 2.5µL of each PCR product and 2.5µL 2X loading buffer with visible dye per sample, per amplification. Let gel run in the dark at approximately 160 volts for 45 minutes or until sample runs far enough to see separate bands for single and double stranded product (bromophenol blue band or other visible marker migrates 1 to 2 inches from wells).

5. Photograph using UV Transilluminator accompanied by a GelStar® Yellow Photographic Filter (Lonza Group, Ltd. No 50536).

CAUTION: Wear protective equipment when handling GelStar® Nucleic Acid Stain or Ethidium Bromide and when photographing gel using UV Transilluminator.

6. Gel analysis

	HLA-DPB	HLA-DQA
Double Strand(s) (bp)	~280	~280, ~240
Single Strand(s) (bp)	~180	~180, ~160

Gel Interpretation

Well

Amplification

Non-amplification

Double Stranded DNA

Single Stranded DNA

Primer Band

-----(Bright)

------(less bright)